## STIC Biotechnology Systems Branch

## RAW SEQUENCE LISTING ERROR REPORT

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<a href="http://www.uspto.gov/ebc/efs/downloads/documents.htm">http://www.uspto.gov/ebc/efs/downloads/documents.htm</a>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
  U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
  Alexandria, VA 22314

Revised 01/10/06



DATE: 02/21/2007

**IFWP** 

PATENT APPLICATION: US/10/524,724 TIME: 14:09:09 Input Set : A:\pto.da.txt Output Set: N:\CRF4\02212007\J524724.raw 3 <110> APPLICANT: DUNCKLEY, et al. 5 <120> TITLE OF INVENTION: Modified Tailed Oligonucleotides 7 <130> FILE REFERENCE: E072 1050.1 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/524,724 11 <141> CURRENT FILING DATE: 2005-02-16 13 <150> PRIOR APPLICATION NUMBER: PCT/GB2003/003612 14 <151> PRIOR FILING DATE: 2003-08-18 16 <150> PRIOR APPLICATION NUMBER: GB 0219143.5 Does Not Comply Corrected Diskette Needed 17 <151> PRIOR FILING DATE: 2002-08-16 19 <160> NUMBER OF SEQ 1D NOS: 16 21 <170> SOFTWARE: PatentIn version 3.1 ERRORED SEQUENCES 23 <210> SEQ ID NO: 1 24 <211> LENGTH: 21 25 <212> TYPE: DNA 26 <213> ORGANISM: Artificial Sequence 28 <220> FEATURE: 29 <223> OTHER INFORMATION: PCR primer 31 <400> SEQUENCE: 1 E--> 32 gggctgctgg ttgtctaccc a --33 21 35 <210> SEQ ID NO: 2 36 <211> LENGTH: 27 37 <212> TYPE: DNA 38 <213> ORGANISM: Artificial sequence 40 <220> FEATURE: 41 <223> OTHER INFORMATION: PCR Primer 43 <400> SEQUENCE: 2 E--> 44 aacttacctg ccaaaaatga tgagaca 45 27 47 <210> SEQ ID NO: 3 48 <211> LENGTH: 41 49 <212> TYPE: DNA 50 <213> ORGANISM: Artificial sequence 52 <220> FEATURE: 53 <223> OTHER INFORMATION: PCR primer 55 <400> SEOUENCE: 3 E--> 56 attcatgtta tatggtcgac agactatcaa cttaatttct g,7 4

RAW SEQUENCE LISTING

59 <210> SEQ ID NO: 4

DATE: 02/21/2007

TIME: 14:09:09

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Input Set : A:\pto.da.txt
                     Output Set: N:\CRF4\02212007\J524724.raw
     60 <211> LENGTH: 35
     61 <212> TYPE: DNA
     62 <213> ORGANISM: Artificial sequence
     64 <220> FEATURE:
     65 <223> OTHER INFORMATION: PCR primer
     67 <400> SEQUENCE: 4
E--> 68 gttagcagag tcgacgtgat ataaaatggc atatc
    69:35
     71 <210> SEQ ID NO: 5
     72 <211> LENGTH: 26
     73 <212> TYPE: DNA
     74 <213> ORGANISM: Artificial sequence
     76 <220> FEATURE:
     77 <223> OTHER INFORMATION: PCR primer
     79 <400> SEQUENCE: 5
E--> 80 cattccttaa atcggagtaa gtctgc
     81 26~
     83 <210> SEQ ID NO: 6
     84_<211> LENGTH: 27
     85 <212> TYPE: DNA
     86 <213> ORGANISM: Artificial sequence
     88 <220> FEATURE:
     89 <223> OTHER INFORMATION: PCR primer
     91 <400> SEQUENCE: 6
E--> 92 gcagacttac tcctgattta aggaatg
     93 27_
     95 <210> SEQ ID NO: 7
     96 <211> LENGTH: 42
     97 <212> TYPE: DNA
     98 <213> ORGANISM: Artificial sequence
     100 <220> FEATURE:
     101 <223> OTHER INFORMATION: PCR primer
     103 <400> SEQUENCE: 7
E--> 104 aaattaatac gactcactat agggctgctg gttgtctacc ca
     105 42 ---
     107 <210> SEQ ID NO: 8
     108 <211> LENGTH: 38
     109 <212> TYPE: RNA
     110 <213> ORGANISM: Artificial sequence
     112 <220> FEATURE:
     113 <223> OTHER INFORMATION: modified tailed oligonucleotide
     115 <220> FEATURE:
     116 <221> NAME/KEY: misc feature
     117 <222> LOCATION: (1)..(23)
     118 <223> OTHER INFORMATION: tail region
     120 <220> FEATURE:
     121 <221> NAME/KEY: misc feature
     122 <222> LOCATION: (24)..(38)
     123 <223> OTHER INFORMATION: complementary region
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/524,724

DATE: 02/21/2007

TIME: 14:09:09

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Input Set : A:\pto.da.txt
                     Output Set: N:\CRF4\02212007\J524724.raw
     125 <400> SEQUENCE: 8
E--> 126 agaagaacga agaacgaaga acagauuuug ucuaaaac
     129 <210> SEQ ID NO: 9
     130 <211> LENGTH: 38
     131 <212> TYPE: RNA
     132 <213> ORGANISM: Artificial sequence. ,
                                    A second contract of the
     134 <220> FEATURE:
     135 <223> OTHER INFORMATION: modified tailed oligonucleotide
     137 <220> FEATURE:
     138 <221> NAME/KEY: misc_feature
     139 <222> LOCATION: (1)..(15)
     140 <223> OTHER INFORMATION: complementary region
     142 <220> FEATURE:
     143 <221> NAME/KEY: misc_feature
     144 <222> LOCATION: (16)..(38)
     145 <223> OTHER INFORMATION: tail region
     147 <400> SEQUENCE: 9
E--> 148 gauuuugucu aaaacagaag aacgaagaac gaagaaca -
     149 38 ---
     151 <210> SEQ ID NO: 10
     152 <211> LENGTH: 38
     153 <212> TYPE: RNA
     154 <213> ORGANISM: Artificial sequence
     156 <220> FEATURE:
     157 <223> OTHER INFORMATION: modified tailed oligonucleotide
     159 <220> FEATURE:
     160 <221> NAME/KEY: misc feature
     161 <222> LOCATION: (1)..(23)
     162 <223> OTHER INFORMATION: tail region
     164 <220> FEATURE:
     165 <221> NAME/KEY: misc feature
     166 <222> LOCATION: (24)..(38)
     167 <223> OTHER INFORMATION: complementary region
     169 <400> SEQUENCE: 10 \
E--> 170 aggaggacgg aggacggagg acagauuuug ucuaaaac
     171 38 -
     173 <210> SEQ ID NO: 11
     174 <211> LENGTH: 38
     175 <212> TYPE: RNA
     176 <213> ORGANISM: Artificial sequence
     178 <220> FEATURE:
     179 <223> OTHER INFORMATION: modified tailed oligonucleotide
     181 <220> FEATURE:
     182 <221> NAME/KEY: misc feature
     183 <222> LOCATION: (1)..(23)
     184 <223> OTHER INFORMATION: tail region
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/524,724

187 <221> NAME/KEY: misc feature

186 <220> FEATURE:

DATE: 02/21/2007

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PATENT APPLICATION: US/10/524,724
                                                              TIME: 14:09:09
                     Input Set : A:\pto.da.txt
                     Output Set: N:\CRF4\02212007\J524724.raw
     188 <222> LOCATION: (24)..(38)
     189 <223> OTHER INFORMATION: complementary region
     191 <400> SEQUENCE: 11
E--> 192 aucuuuccuc uuuccucuuu ccagauuuug ucuaaaac > 3
     193 38 __
     195 <210> SEQ ID NO: 12
     196 <211> LENGTH: 38
     197 <212> TYPE: RNA
     198 <213> ORGANISM: Artificial sequence
     200 <220> FEATURE:
     201 <223> OTHER INFORMATION: modified tailed oligonucleotide
     203 <220> FEATURE:
     204 <221> NAME/KEY: misc feature
     205 <222> LOCATION: (1)..(23)
     206 <223> OTHER INFORMATION: tail region
     208 <220> FEATURE:
     209 <221> NAME/KEY: misc_feature
     210 <222> LOCATION: (24)..(38)
     211 <223> OTHER INFORMATION: complementary region.
                                                               and the second second
     213 <400> SEQUENCE: 12
E--> 214 auagggcagg cuagggcagg ccagauuuug ucuaaaac
     215 38 ~
     217 <210> SEQ ID NO: 13
     218 <211> LENGTH: 15
     219 <212> TYPE: RNA
     220 <213> ORGANISM: Artificial sequence
     222 <220> FEATURE:
     223 <223> OTHER INFORMATION: modified tailed oligonucleotide
    '225 <220> FEATURE:
     226 <221> NAME/KEY: misc feature
     227 <222> LOCATION: (1)..(15)
     228 <223> OTHER INFORMATION: complementary region
     230 <400> SEQUENCE: 13
E--> 231 gauuuugucu aaaac
     232 15 -
     234 <210> SEQ ID NO: 14
     235 <211> LENGTH: 14
     236 <212> TYPE: RNA
     237 <213> ORGANISM: Artificial sequence
     239 <220> FEATURE:
     240 <223> OTHER INFORMATION: modified tailed oligonucleotide
     242 <220> FEATURE:
     243 <221> NAME/KEY: misc_feature
     244 <222> LOCATION: (1)..(14)
     245 <223> OTHER INFORMATION: complementary region
     247 <400> SEQUENCE: 14
E--> 248 acccugucuu aggu
     249 14 ___
     272 <210> SEQ ID NO: 16
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RAW SEQUENCE LISTING

RAW SEQUENCE LISTING

DATE: 02/21/2007

PATENT APPLICATION: US/10/524,724

TIME: 14:09:09

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\02212007\J524724.raw

274 <211> LENGTH: 23

275 <212> TYPE: RNA

276 <213> ORGANISM: Artificial sequence

278 <220> FEATURE:

279 <223> OTHER INFORMATION: tail region for modified tailed oligonucleotide

281 <400> SEQUENCE: 16

E--> 282 aggaggacgg aggacggagg aca 7 2

283 23

E--> 285 1 E--> 288 1

PIs delete

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/524,724

DATE: 02/21/2007 TIME: 14:09:10

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\02212007\J524724.raw

## Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 32 Seq#:2; Line(s) 44 Seq#:3; Line(s) 56 Seg#:4; Line(s) 68 Seq#:5; Line(s) 80 Seq#:6; Line(s) 92 Seq#:7; Line(s) 104 Seq#:8; Line(s) 126 Seq#:9; Line(s) 148 Seq#:10; Line(s) 170 Seq#:11; Line(s) 192 Seq#:12; Line(s) 214 Seq#:13; Line(s) 231 Seq#:14; Line(s) 248 Seq#:15; Line(s) 270 Seq#:16; Line(s) 282

## VERIFICATION SUMMARY

DATE: 02/21/2007 PATENT APPLICATION: US/10/524,724 TIME: 14:09:10

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\02212007\J524724.raw

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L:32 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:21 SEQ:1
L:44 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:27 SEQ:2
L:56 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:41 SEQ:3
L:68 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:35 SEQ:4
L:80 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:26 SEQ:5
L:92 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:27 SEQ:6
L:104 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:42 SEQ:7 L:126 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:38 SEQ:8
L:148 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:38 SEQ:9
L:170 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:38 SEQ:10
L:192 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:38 SEQ:11
L:214 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:38 SEQ:12
L:231 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:15 SEQ:13
L:248 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:14 SEQ:14 L:282 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:23 SEQ:16
L:285 M:254 E: No. of Bases conflict, this line has no nucleotides.
L:288 M:254 E: No. of Bases conflict, this line has no nucleotides.
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